```
<110>
           Shaughnessy, John D.
          Evi27 Gene Sequence and Protein Encoded Thereby
<120>
<130>
          D6138
<140>
          US 09/778,971
<141>
           2001-02-02
<150>
          US 60/180,374
<151>
           2000-02-04
           9
<160>
<210>
           1827
<211>
<212>
          DNA
<213>
          Homo sapiens
<220>
<221>
          prim_transcript
<223>
          cDNA of human Evi27
<400>
           1
```

50 cggcgatgtc gctcgtgctg ataagcctgg ccgcgctgtg caggagcgcc gtaccccgag agccgaccgt tcaatgtggc tctgaaactg ggccatctcc 100 150 agagtggatg ctacaacatg atctaatccc cggagacttg agggacctcc gagtagaacc tgttacaact agtgttgcaa caggggacta ttcaattttg 200 atqaatqtaa qctqqqtact ccqqqcaqat qccaqcatcc qcttgttgaa 250 ggccaccaag atttgtgtga cgggcaaaag caacttccag tcctacagct 300 350 gtgtgaggtg caattacaca gaggccttcc agactcagac cagaccctct qqtqqtaaat ggacattttc ctatatcggc ttccctgtag agctgaacac 400 450 agtctatttc attggggccc ataatattcc taatgcaaat atgaatgaag 500 atggecette catgtetgtg aattteacet caccaggetg cetagaceae 550 ataatgaaat ataaaaaaa gtgtgtcaag gccggaagcc tgtgggatcc 600 qaacatcact gcttgtaaga agaatgagga gacagtagaa gtgaacttca caaccactcc cctgggaaac agatacatgg ctcttatcca acacagcact 650 atcatcgggt tttctcaggt gtttgagcca caccagaaga aacaaacgcg 700 agetteagtg gtgatteeag tgaetgggga tagtgaaggt getaeggtge 750 800 agetgaetee atatttteet aettgtggea gegaetgeat eegaeataaa 850 ggaacagttg tgctctgccc acaaacaggc gtccctttcc ctctggataa caacaaaagc aagccgggag gctggctgcc tctcctcctg ctgtctctgc 900 950 tggtggcac atgggtgctg gtggcaggga tctatctaat gtggaggcac gaaaggatca agaagacttc cttttctacc accacactac tgcccccat 1000 taaggttett gtggtttacc catetgaaat atgtttecat cacacaattt 1050 gttacttcac tgaatttctt caaaaccatt gcagaagtga ggtcatcctt 1100 gaaaagtggc agaaaaagaa aatagcagag atgggtccag tgcagtggct 1150 tgccactcaa aagaaggcag cagacaaagt cgtcttcctt ctttccaatg 1200 acgtcaacag tgtgtgcgat ggtacctgtg gcaagagcga gggcagtccc 1250

```
agtgagaact ctcaagacct cttccccctt gcctttaacc ttttctgcag 1300
tgatctaaga agccagattc atctgcacaa atacgtggtg gtctacttta 1350
gagagattga tacaaaagac gattacaatg ctctcagtgt ctgccccaag 1400
taccaettea tgaaggatge caetgettte tgtgeagaae ttetecatgt 1450
caagcagcag gtgtcagcag gaaaaagatc acaagcctgc cacgatggct 1500
gctgctcctt gtagcccacc catgagaagc aagagacctt aaaggcttcc 1550
tatcccacca attacaggga aaaaacgtgt gatgatcctg aagcttacta 1600
tgcagcctac aaacagcctt agtaattaaa acattttata ccaataaaat 1650
tttcaaatat tactaactaa tgtagcatta actaacgatt ggaaactaca 1700
tttacaactt caaagctgtt ttatacatag aaatcaatta cagctttaat 1750
tgaaaactgt aaccattttg ataatgcaac aataaagcat cttccaaaaa 1800
aaaaaaaaa aaaaaaaa aaaaaaa
                                                        1827
     <210>
                2
               2856
     <211>
     <212>
               DNA
     <213>
               Homo sapiens
     <220>
               prim_transcript
     <221>
     <223>
               cDNA of human Evi27
     <400>
                                                          50
eggegatgte getegtgetg ataageetgg eegegetgtg eaggagegee
                                                         100
qtaccccqaq aqccqaccqt tcaatqtqqc tctqaaactq ggccatctcc
                                                         150
agagtggatg ctacaacatg atctaatccc cggagacttg agggacctcc
                                                         200
qaqtaqaacc tgttacaact agtgttgcaa caggggacta ttcaattttg
atgaatgtaa gctgggtact ccgggcagat gccagcatcc gcttgttgaa
                                                         250
ggccaccaag atttgtgtga cgggcaaaag caacttccag tcctacagct
                                                         300
                                                         350
qtqtgaggtg caattacaca gaggccttcc agactcagac cagaccctct
                                                         400
ggtggtaaat ggacattttc ctatatcggc ttccctgtag agctgaacac
                                                         450
agtctatttc attggggccc ataatattcc taatgcaaat atgaatgaag
atggcccttc catgtctgtg aatttcacct caccaggctg cctagaccac
                                                         500
ataatgaaat ataaaaaaa gtgtgtcaag gccggaagcc tgtgggatcc
                                                         550
                                                        600
qaacatcact qcttqtaaqa agaatgagga gacagtagaa gtgaacttca
                                                         650
caaccactcc cctgggaaac agatacatgg ctcttatcca acacagcact
                                                         700
atcatcgggt tttctcaggt gtttgagcca caccagaaga aacaaacgcg
                                                         750
agetteagtg gtgatteeag tgactgggga tagtgaaggt getaeggtge
aggtaaagtt cagtgagctg ctctggggag ggaagggaca tagaagactg
                                                         800
ttccatcatt cattgctttt aaggatgagt tctctcttgt caaatgcact
                                                         850
                                                         900
tetgecagea gacaccagtt aagtggegtt catgggggtt etttegetge
                                                         950
agectecace gtgctgaggt caggaggeeg acgtggcagt tgtggteect
tttgcttgta ttaatggctg ctgaccttcc aaagcacttt ttattttcat 1000
tttctgtcac agacactcag ggatagcagt accattttac ttccgcaagc 1050
ctttaactgc aagatgaagc tgcaaagggt ttgaaatggg aaggtttgag 1100
ttccaggcag cgtatgaact ctggagaggg gctgccagtc ctctctgggc 1150
cgcagcggac ccagctggaa cacaggaagt tggagcagta ggtgctcctt 1200
caccteteag tatgtetett teaactetag tttttgaagt ggggacaeag 1250
gaagtccagt ggggacacag ccactcccca aagaataagg aacttccatg 1300
cttcattccc tggcataaaa agtgntcaaa cacaccagag ggggcaggca 1350
```

ccagccaggg tatgatgggt actaccettt tetggagaac catagaette 1400

```
ccttactaca gggacttgca tgtcctaaag cactggctga aggaagccaa 1450
gaggatcact gctgctcctt ttttgtagag gaaatgtttg tgtacgtggt 1500
aagatatgac ctagcccttt taggtaagcg aactggtatg ttagtaacgt 1550
gtacaaagtt taggttcaga ccccgggagt cttgggcatg tgggtctcgg 1600
gtcactggtt ttgactttag ggctttgtta cagatgtgtg accaagggga 1650
aaatgtgcat gacaacacta gaggtagggg cgaagccaga aagaagggaa 1700
gttttggctg aagtaggagt cttggtgaga ttttgctgtg atgcatggtg 1750
tgaactttet gageetettg ttttteetea getgaeteea tatttteeta 1800
cttgtggcag cgactgcatc cgacataaag gaacagttgt gctctgccca 1850
caaacaggcg tccctttccc tctggataac aacaaaagca agccgggagg 1900
etggetgeet eteeteetge tgtetetget ggtggeeaea tgggtgetgg 1950
tggcagggat ctatctaatg tggaggcacg aaaggatcaa gaagacttcc 2000
ttttctacca ccacactact gcccccatt aaggttcttg tggtttaccc 2050
atctgaaata tgtttccatc acacaatttg ttacttcact gaatttcttc 2100
aaaaccattg cagaagtgag gtcatccttg aaaagtggca gaaaaagaaa 2150
atagcagaga tgggtccagt gcagtggctt gccactcaaa agaaggcagc 2200
agacaaagtc gtcttccttc tttccaatga cgtcaacagt gtgtgcgatg 2250
gtacctgtgg caagagcgag ggcagtccca gtgagaactc tcaagacctc 2300
ttcccccttg cctttaacct tttctgcagt gatctaagaa gccagattca 2350
tctgcacaaa tacgtggtgg tctactttag agagattgat acaaaagacg 2400
attacaatgc tctcagtgtc tgccccaagt accacttcat gaaggatgcc 2450
actgetttet gtgeagaact tetecatgte aageageagg tgteageagg 2500
aaaaagatca caagcctgcc acgatggctg ctgctccttg tagcccaccc 2550
atgagaagca agagacctta aaggcttcct atcccaccaa ttacagggaa 2600
aaaacgtgtg atgateetga agettaetat geageetaea aacageetta 2650
gtagcattaa ctaacgattg gaaactacat ttacaacttc aaagctgttt 2750
tatacataga aatcaattac agctttaatt gaaaactgta accattttga 2800
aaaaaa
                                                     2856
    <210>
              1963
    <211>
    <212>
              DNA
    <213>
              Unknown
    <220>
    <221>
              prim_transcript
    <223>
              cDNA of mouse Evi27
    <400>
qtqqccaqtq gccgggccat gttgctagtq ttgctgatct tggctgcatc
                                                       50
gtgcaggagc gccctgcctc gagagccgac tattcagtgt ggctctgaga
                                                      100
cagggccatc tccagagtgg atggtccaac acacactcac tccaggagac
                                                      150
ttgagggacc tccaagtgga actcgtcaag acaagtgtgg cagcagagga
                                                      200
gttttcaatt ttgatgaaca taagctggat actccgggca gacgccagca
                                                      250
tccgcttgtt gaaggccacc aagatctgcg tgagtggcaa aaacaacatg
                                                      300
aattcataca gctgtgtgag gtgcaactac acagaggcct tccaaagcca
                                                      350
gaccagacct teeggeggea aatggacatt eteetatgta ggetteeetg
                                                      400
tggagetgag cactetetat etcateageg cecataacat ceceaatget
                                                      450
aatatgaatg aggacagccc ttctttgtct gtgaacttca cctcgccagg
                                                      500
```

```
550
ctgcctaaac cacgtaatga aatataaaaa gcagtgcact gaggcgggaa
                                                         600
gcctgtggga cccagacatc actgcttgta aaaagaacga gaagatggtt
                                                         650
qaaqtqaatt tcacaaccaa tccccttgga aacagataca cgattctcat
tcaacgggac acgacattgg ggttttctag agtgctggag aataaactga
                                                         700
                                                         750
tgaggacgtc tgtagccatc ccggtgactg aggagagtga aggtgcggtg
gttcagctga ccccatattt acatacctgc ggcaatgact gcatccgacg
                                                         800
cqaagggaca gttgtgcttt gctcagagac aagtgctccc atccctccag
                                                         850
atgacaacag acgcatgctg ggaggctggc tgcctctctt cctggtgctg
                                                         900
                                                         950
ctggtggctg tgtgggtgct ggcagctggg atctacctaa cttggaggca
aggaaggagc acgaagacgt cetttectat ttecaccatg etectgeece 1000
tcattaaggt cctggtggtt tatccttctg agatatgttt ccatcacacc 1050
gtctgtcgct tcactgactt tcttcaaaac tactgcagaa gtgaggtcat 1100
ccttgaaaaa tggcagaaaa agaaaatcgc cgagatgggg ccggtacagt 1150
ggctgaccac tcagaagcaa gcggcagata aagtggtctt ccttcttccc 1200
agtgacgtcc cgaccctttg tgacagtgcc tgtggccaca atgagggcag 1250
cgccagggag aactctcagg atctgttccc tcttgccttt aacctctttt 1300
gtagtgattt cagcagccag acgcatctgc acaaatacct ggtggtctat 1350
cttgggggag cagacctcaa aggcgactat aatgccctga gtgtctgccc 1400
ccaatatcat ctcatgaagg acgccacagc tttccacaca gaacttctca 1450
aggetaegea gageatgtea gtgaagaaac geteacaage etgeeatgat 1500
agetgtteac cettgtagte caceeggggg aatagagaet etgaageett 1550
cctactctcc cttccagtga caaatgctgt gtgacgactc tgaaatgtgt 1600
gggagagget gtgtggaggt agtgctatgt acaaacttgc tttaaaactg 1650
gagtttgcaa agtcaacctg agcatacacg cctgaggcta gtcattggct 1700
ggatttatga agacaacaca gttacagaca ataatgagtg ggacctacat 1750
ttgggatata cccaaagctg ggtaatgatt atcactgaga accacgcact 1800
ctggccatga agtaatacgg cacttccctg tcaggctgtc tgtcaggttg 1850
ggtctgtctt gcactgccca tgctctatgc tgcacgtaga ccgttttgta 1900
acattttaat ctgttaatga ataatccgtt tgggaagctc tcaaaaaaaa 1950
                                                        1963
aaaaaaaaa aaa
     <210>
               2589
     <211>
     <212>
               DNA
     <213>
               Unknown
     <220>
     <221>
               prim_transcript
     <223>
               cDNA of mouse Evi27
     <400>
                                                          50
gtggccagtg gccgggccat gttgctagtg ttgctgatct tggctgcatc
                                                         100
gtgcaggagc gccctgcctc gagagccgac tattcagtgt ggctctgaga
                                                         150
cagggccatc tccagagtgg atggtccaac acacactcac tccaggagac
ttgagggacc tccaagtgga actcgtcaag acaagtgtgg cagcagagga
                                                         200
gttttcaatt ttgatgaaca taagctggat actccgggca gacgccagca
                                                         250
                                                        300
tccgcttgtt gaaggccacc aagatctgcg tgagtggcaa aaacaacatg
                                                         350
aattcataca gctgtgtgag gtgcaactac acagaggcct tccaaagcca
                                                         400
qaccaqacct teeggeggea aatggacatt eteetatgta ggetteeetg
                                                         450
tggagctgag cactctctat ctcatcagcg cccataacat ccccaatgct
aatatgaatg aggacagccc ttctttgtct gtgaacttca cctcgccagg
                                                         500
```

```
550
qtqcactcgt gaaaacacag aagtaacgtc cggtgtattt ccagcagcta
aacaccaggc teteeggatt teageteett teecattaca attteeteet
                                                         600
                                                         650
gggccagagg actcagtcat tetgccacec cageetetgg egtegetttt
tcatgacttt gtcaaactta cctagcttgt ttccattctg aaattgtctg
                                                        700
atgcttgctt cgtatgtaag ccggggatat gaggtttggg tatgaatccc
                                                        750
acagagggca ctgaattctt ctcactatgg cctatctggg ctgtgtgaca
                                                        800
                                                         850
ttgttggtga gggtcgtgcc tactaggcat ctgggtatct accacctgga
gcttcatgtc tggaagaggc agaacctata tgtattgtca gctctcactt
                                                         900
                                                         950
ttgtttccgt gtcacctcct ggagactgtt tttgataaag gttgtactta
aaggagatta cttaaagctt ccgtggaaga atggtttcct atttagatct 1000
gttgtctctc atatctgaag taagtgtgtg tgtgtgtgtt ttgtgtgtgt 1050
gtgtgtgtgt gtgtgtatac tgggcaaagg gttatacctt tactcaaatg 1100
taacaacttt cattcacatt cccaggctgc ctaaaccacg taatgaaata 1150
taaaaagcag tgcactgagg cgggaagcct gtgggaccca gacatcactg 1200
cttgtaaaaa gaacgagaag atggttgaag tgaatttcac aaccaatccc 1250
cttggaaaca gatacacgat tctcattcaa cgggacacga cattggggtt 1300
ttctagagtg ctggagaata aactgatgag gacgtctgta gccatcccgg 1350
tgactgagga gagtgaaggt gcggtggttc agctgacccc atatttacat 1400
acctgcggca atgactgcat ccgacgcgaa gggacagttg tgctttgctc 1450
agagacaagt gctcccatcc ctccagatga caacagacgc atgctgggag 1500
gctggctgcc tctcttcctg gtgctgctgg tggctgtgtg ggtgctggca 1550
gctgggatct acctaacttg gaggcaagga aggagcacga agacgtcctt 1600
tectatttee accatgetee tgeeceteat taaggteetg gtggtttate 1650
cttctgagat atgtttccat cacaccgtct gtcgcttcac tgactttctt 1700
caaaactact gcagaagtga ggtcatcctt gaaaaaatggc agaaaaagaa 1750
aatcgccgag atggggccgg tacagtggct gaccactcag aagcaagcgg 1800
cagataaagt ggtcttcctt cttcccagtg acgtcccgac cctttgtgac 1850
agtgcctgtg gccacaatga gggcagcgcc agggagaact ctcaggatct 1900
gttccctctt gcctttaacc tcttttgtag tgatttcagc agccagacgc 1950
atctgcacaa atacctggtg gtctatcttg ggggagcaga cctcaaaggc 2000
gactataatg ccctgagtgt ctgcccccaa tatcatctca tgaaggacgc 2050
cacagettte cacacagaac tteteaagge tacgeagage atgteagtga 2100
agaaacgctc acaagcctgc catgatagct gttcaccctt gtagtccacc 2150
cgggggaata gagactetga ageetteeta eteteeette cagtgacaaa 2200
tgctgtgtga cgactctgaa atgtgtggga gaggctgtgt ggaggtagtg 2250
ctatgtacaa acttgcttta aaactggagt ttgcaaagtc aacctgagca 2300
tacacgcctg aggctagtca ttggctggat ttatgaagac aacacagtta 2350
cagacaataa tgagtgggac ctacatttgg gatataccca aagctgggta 2400
atgattatca ctgagaacca cgcactctgg ccatgaagta atacggcact 2450
tecetgteag getgtetgte aggttgggte tgtettgeae tgeecatget 2500
ctatgctgca cgtagaccgt tttgtaacat tttaatctgt taatgaataa 2550
                                                       2589
tccgtttggg aagctctcaa aaaaaaaaa aaaaaaaaa
               5
     <210>
               502
     <211>
     <212>
               PRT
     <213>
               Homo sapiens
     <220>
     <221>
               peptide
     <223>
               Human Evi27 protein
```

<400> Met Ser Leu Val Leu Ile Ser Leu Ala Ala Leu Cys Arg Ser Ala Val Pro Arg Glu Pro Thr Val Gln Cys Gly Ser Glu Thr Gly Pro Ser Pro Glu Trp Met Leu Gln His Asp Leu Ile Pro Gly Asp Leu Arg Asp Leu Arg Val Glu Pro Val Thr Thr Ser Val Ala Thr Gly Asp Tyr Ser Ile Leu Met Asn Val Ser Trp Val Leu Arg Ala Asp Ala Ser Ile Arg Leu Leu Lys Ala Thr Lys Ile Cys Val Thr Gly Lys Ser Asn Phe Gln Ser Tyr Ser Cys Val Arg Cys Asn Tyr Thr Glu Ala Phe Gln Thr Gln Thr Arg Pro Ser Gly Gly Lys Trp Thr Phe Ser Tyr Ile Gly Phe Pro Val Glu Leu Asn Thr Val Tyr Phe Ile Gly Ala His Asn Ile Pro Asn Ala Asn Met Asn Glu Asp Gly Pro Ser Met Ser Val Asn Phe Thr Ser Pro Gly Cys Leu Asp His Ile Met Lys Tyr Lys Lys Cys Val Lys Ala Gly Ser Leu Trp Asp Pro Asn Ile Thr Ala Cys Lys Lys Asn Glu Glu Thr Val Glu Val Asn Phe Thr Thr Pro Leu Gly Asn Arg Tyr Met Ala Leu Ile Gln His Ser Thr Ile Ile Gly Phe Ser Gln Val Phe Glu Pro His Gln Lys Lys Gln Thr Arg Ala Ser Val Val Ile Pro Val Thr Gly Asp Ser Glu Gly Ala Thr Val Gln Leu Thr Pro Tyr Phe Pro Thr Cys Gly Ser Asp Cys Ile Arg His Lys Gly Thr Val Val Leu Cys Pro Gln Thr Gly Val Pro Phe Pro Leu Asp Asn Asn Lys Ser Lys Pro Gly Gly Trp Leu Pro Leu Leu Leu Ser Leu Leu Val Ala Thr Trp Val Leu Val Ala Gly Ile Tyr Leu Met Trp Arg His Glu Arg Ile Lys Lys Thr Ser Phe Ser Thr Thr Thr Leu Leu Pro Pro Ile Lys Val Leu Val Val Tyr Pro Ser Glu Ile Cys Phe His His Thr Ile Cys Tyr Phe Thr Glu Phe Leu Gln Asn His Cys Arg Ser Glu Val Ile Leu Glu Lys Trp Gln Lys Lys Lys Ile Ala Glu

```
Met Gly Pro Val Gln Trp Leu Ala Thr Gln Lys Lys Ala Ala Asp
                 380
                                      385
Lys Val Val Phe Leu Leu Ser Asn Asp Val Asn Ser Val Cys Asp
                 395
                                     400
Gly Thr Cys Gly Lys Ser Glu Gly Ser Pro Ser Glu Asn Ser Gln
                410
                                     415
Asp Leu Phe Pro Leu Ala Phe Asn Leu Phe Cys Ser Asp Leu Arg
                 425
                                      430
Ser Gln Ile His Leu His Lys Tyr Val Val Val Tyr Phe Arg Glu
                 440
                                     445
                                                          450
Ile Asp Thr Lys Asp Asp Tyr Asn Ala Leu Ser Val Cys Pro Lys
                 455
                                     460
Tyr His Phe Met Lys Asp Ala Thr Ala Phe Cys Ala Glu Leu Leu
                 470
                                     475
His Val Lys Gln Gln Val Ser Ala Gly Lys Arg Ser Gln Ala Cys
                                      490
                                                          495
                 485
His Asp Gly Cys Cys Ser Leu
                500
     <210>
                6
     <211>
                288
     <212>
                PRT
     <213>
                Homo sapiens
     <220>
                peptide
     <221>
     <223>
                Human Evi27 protein
     <400>
Met Ser Leu Val Leu Ile Ser Leu Ala Ala Leu Cys Arg Ser Ala
Val Pro Arg Glu Pro Thr Val Gln Cys Gly Ser Glu Thr Gly Pro
                 20
                                      25
Ser Pro Glu Trp Met Leu Gln His Asp Leu Ile Pro Gly Asp Leu
                  35
                                      40
Arg Asp Leu Arg Val Glu Pro Val Thr Thr Ser Val Ala Thr Gly
                                      55
                  50
Asp Tyr Ser Ile Leu Met Asn Val Ser Trp Val Leu Arg Ala Asp
Ala Ser Ile Arg Leu Leu Lys Ala Thr Lys Ile Cys Val Thr Gly
                                                           90
                  80
                                      85
Lys Ser Asn Phe Gln Ser Tyr Ser Cys Val Arg Cys Asn Tyr Thr
                                                          105
                  95
                                     100
Glu Ala Phe Gln Thr Gln Thr Arg Pro Ser Gly Gly Lys Trp Thr
                 110
                                     115
                                                          120
Phe Ser Tyr Ile Gly Phe Pro Val Glu Leu Asn Thr Val Tyr Phe
                                                          135
                 125
                                      130
Ile Gly Ala His Asn Ile Pro Asn Ala Asn Met Asn Glu Asp Gly
                 140
                                     145
Pro Ser Met Ser Val Asn Phe Thr Ser Pro Gly Cys Leu Asp His
                 155
                                     160
                                                          165
```

```
Ile Met Lys Tyr Lys Lys Cys Val Lys Ala Gly Ser Leu Trp
                170
Asp Pro Asn Ile Thr Ala Cys Lys Lys Asn Glu Glu Thr Val Glu
                185
                                     190
Val Asn Phe Thr Thr Pro Leu Gly Asn Arg Tyr Met Ala Leu
                                     205
                200
Ile Gln His Ser Thr Ile Ile Gly Phe Ser Gln Val Phe Glu Pro
                215
                                     220
His Gln Lys Lys Gln Thr Arg Ala Ser Val Val Ile Pro Val Thr
                230
                                     235
                                                          240
Gly Asp Ser Glu Gly Ala Thr Val Gln Val Lys Phe Ser Glu Leu
                245
                                     250
Leu Trp Gly Gly Lys Gly His Arg Arg Leu Phe His His Ser Leu
                260
                                     265
                                                          270
Leu Leu Arg Met Ser Ser Leu Leu Ser Asn Ala Leu Leu Pro Ala
                275
                                     280
                                                          285
Asp Thr Ser
        288
     <210>
                7
                499
     <211>
     <212>
                PRT
     <213>
               Unknown
     <220>
     <221>
               peptide
               Mouse Evi27 protein
     <223>
     <400>
Met Leu Val Leu Leu Ile Leu Ala Ala Ser Cys Arg Ser Ala
Leu Pro Arg Glu Pro Thr Ile Gln Cys Gly Ser Glu Thr Gly Pro
                 20
Ser Pro Glu Trp Met Val Gln His Thr Leu Thr Pro Gly Asp Leu
                                      40
                 35
Arg Asp Leu Gln Val Glu Leu Val Lys Thr Ser Val Ala Ala Glu
                 50
                                      55
Glu Phe Ser Ile Leu Met Asn Ile Ser Trp Ile Leu Arg Ala Asp
Ala Ser Ile Arg Leu Leu Lys Ala Thr Lys Ile Cys Val Ser Gly
                                                           90
                                      85
                 80
Lys Asn Asn Met Asn Ser Tyr Ser Cys Val Arg Cys Asn Tyr Thr
                                     100
                                                          105
                 95
Glu Ala Phe Gln Ser Gln Thr Arg Pro Ser Gly Gly Lys Trp Thr
                110
                                     115
                                                          120
Phe Ser Tyr Val Gly Phe Pro Val Glu Leu Ser Thr Leu Tyr Leu
                                     130
                                                          135
                125
Ile Ser Ala His Asn Ile Pro Asn Ala Asn Met Asn Glu Asp Ser
                140
                                     145
                                                          150
Pro Ser Leu Ser Val Asn Phe Thr Ser Pro Gly Cys Leu Asn His
                155
                                     160
                                                          165
```

```
Val Met Lys Tyr Lys Lys Gln Cys Thr Glu Ala Gly Ser Leu Trp
                170
                                     175
Asp Pro Asp Ile Thr Ala Cys Lys Lys Asn Glu Lys Met Val Glu
                185
                                     190
Val Asn Phe Thr Thr Asn Pro Leu Gly Asn Arg Tyr Thr Ile Leu
                200
                                     205
Ile Gln Arg Asp Thr Thr Leu Gly Phe Ser Arg Val Leu Glu Asn
                215
                                     220
Lys Leu Met Arg Thr Ser Val Ala Ile Pro Val Thr Glu Glu Ser
                230
                                     235
                                                          240
Glu Gly Ala Val Val Gln Leu Thr Pro Tyr Leu His Thr Cys Gly
                245
                                     250
Asn Asp Cys Ile Arg Arg Glu Gly Thr Val Val Leu Cys Ser Glu
                260
                                     265
                                                          270
Thr Ser Ala Pro Ile Pro Pro Asp Asp Asn Arg Arg Met Leu Gly
                275
                                     280
                                                          285
Gly Trp Leu Pro Leu Phe Leu Val Leu Leu Val Ala Val Trp Val
                290
                                     295
Leu Ala Ala Gly Ile Tyr Leu Thr Trp Arg Gln Gly Arg Ser Thr
                305
                                     310
Lys Thr Ser Phe Pro Ile Ser Thr Met Leu Leu Pro Leu Ile Lys
                320
                                     325
Val Leu Val Val Tyr Pro Ser Glu Ile Cys Phe His His Thr Val
                335
Cys Arg Phe Thr Asp Phe Leu Gln Asn Tyr Cys Arg Ser Glu Val
                                     355
                350
Ile Leu Glu Lys Trp Gln Lys Lys Ile Ala Glu Met Gly Pro
                365
                                     370
                                                          375
Val Gln Trp Leu Thr Thr Gln Lys Gln Ala Ala Asp Lys Val Val
                                                          390
                380
                                     385
Phe Leu Leu Pro Ser Asp Val Pro Thr Leu Cys Asp Ser Ala Cys
                                                          405
                395
                                     400
Gly His Asn Glu Gly Ser Ala Arg Glu Asn Ser Gln Asp Leu Phe
                410
                                     415
Pro Leu Ala Phe Asn Leu Phe Cys Ser Asp Phe Ser Ser Gln Thr
                425
                                     430
His Leu His Lys Tyr Leu Val Val Tyr Leu Gly Gly Ala Asp Leu
                440
                                     445
Lys Gly Asp Tyr Asn Ala Leu Ser Val Cys Pro Gln Tyr His Leu
                455
Met Lys Asp Ala Thr Ala Phe His Thr Glu Leu Leu Lys Ala Thr
                470
                                     475
Gln Ser Met Ser Val Lys Lys Arg Ser Gln Ala Cys His Asp Ser
                485
                                     490
Cys Ser Pro Leu
            499
     <210>
     <211>
                289
     <212>
                PRT
               Unknown
     <213>
```

```
peptide
     <221>
     <223>
               Mouse Evi27 protein
     <400>
Met Leu Val Leu Leu Ile Leu Ala Ala Ser Cys Arg Ser Ala
Leu Pro Arg Glu Pro Thr Ile Gln Cys Gly Ser Glu Thr Gly Pro
                 20
Ser Pro Glu Trp Met Val Gln His Thr Leu Thr Pro Gly Asp Leu
                 35
                                      40
Arg Asp Leu Gln Val Glu Leu Val Lys Thr Ser Val Ala Ala Glu
                                      55
                 50
Glu Phe Ser Ile Leu Met Asn Ile Ser Trp Ile Leu Arg Ala Asp
Ala Ser Ile Arg Leu Leu Lys Ala Thr Lys Ile Cys Val Ser Gly
                                      85
                 80
Lys Asn Asn Met Asn Ser Tyr Ser Cys Val Arg Cys Asn Tyr Thr
                                     100
                 95
Glu Ala Phe Gln Ser Gln Thr Arg Pro Ser Gly Gly Lys Trp Thr
                                     115
                110
Phe Ser Tyr Val Gly Phe Pro Val Glu Leu Ser Thr Leu Tyr Leu
                                     130
                125
Ile Ser Ala His Asn Ile Pro Asn Ala Asn Met Asn Glu Asp Ser
                140
                                     145
Pro Ser Leu Ser Val Asn Phe Thr Ser Pro Gly Cys Leu Asn His
                155
                                     160
Val Met Lys Tyr Lys Lys Gln Cys Thr Glu Ala Gly Ser Leu Trp
                170
                                     175
Asp Pro Asp Ile Thr Ala Cys Lys Lys Asn Glu Lys Met Val Glu
                185
                                     190
Val Asn Phe Thr Thr Asn Pro Leu Gly Asn Arg Tyr Thr Ile Leu
                200
                                     205
Ile Gln Arg Asp Thr Thr Leu Gly Phe Ser Arg Val Leu Glu Asn
                215
                                     220
Lys Leu Met Arg Thr Ser Val Ala Ile Pro Val Thr Glu Glu Ser
                230
                                     235
Glu Gly Ala Val Val Gln Leu Thr Pro Tyr Leu His Thr Cys Gly
                245
Asn Asp Cys Ile Arg Arg Glu Gly Thr Val Val Leu Cys Ser Glu
                260
                                     265
Thr Ser Ala Pro Ile Pro Pro Asp Asp Asn Arg Arg Met Leu Gly
                275
                                     280
                                                          285
Gly Trp Leu Pro
            289
     <210>
                866
     <211>
     <212>
                PRT
     <213>
               Unknown
```

<220>

<220>

<221> peptide IL-17 receptor protein <400> Met Gly Ala Ala Arg Ser Pro Pro Ser Ala Val Pro Gly Pro Leu Leu Gly Leu Leu Leu Leu Gly Val Leu Ala Pro Gly Gly Ala Ser Leu Arg Leu Leu Asp His Arg Ala Leu Val Cys Ser Gln Pro Gly Leu Asn Cys Thr Val Lys Asn Ser Thr Cys Leu Asp Asp 55 Ser Trp Ile His Pro Arg Asn Leu Thr Pro Ser Ser Pro Lys Asp Leu Gln Ile Gln Leu His Phe Ala His Thr Gln Gln Gly Asp Leu Phe Pro Val Ala His Ile Glu Trp Thr Leu Gln Thr Asp Ala Ser 100 Ile Leu Tyr Leu Glu Gly Ala Glu Leu Ser Val Leu Gln Leu Asn 110 115 Thr Asn Glu Arg Leu Cys Val Arg Phe Glu Phe Leu Ser Lys Leu 125 130 Arg His His Arg Arg Trp Arg Phe Thr Phe Ser His Phe Val 140 145 Val Asp Pro Asp Gln Glu Tyr Glu Val Thr Val His His Leu Pro 155 160 Lys Pro Ile Pro Asp Gly Asp Pro Asn His Gln Ser Lys Asn Phe 170 175 Leu Val Pro Asp Cys Glu His Ala Arg Met Lys Val Thr Thr Pro 185 190 Cys Met Ser Ser Gly Ser Leu Trp Asp Pro Asn Ile Thr Val Glu 200 205 Thr Leu Glu Ala His Gln Leu Arg Val Ser Phe Thr Leu Trp Asn 215 220 Glu Ser Thr His Tyr Gln Ile Leu Leu Thr Ser Phe Pro His Met 230 235 Glu Asn His Ser Cys Phe Glu His Met His His Ile Pro Ala Pro 245 250 Arg Pro Glu Glu Phe His Gln Arg Ser Asn Val Thr Leu Thr Leu 260 265 Arg Asn Leu Lys Gly Cys Cys Arg His Gln Val Gln Ile Gln Pro 275 280 Phe Phe Ser Ser Cys Leu Asn Asp Cys Leu Arg His Ser Ala Thr 290 295 Val Ser Cys Pro Glu Met Pro Asp Thr Pro Glu Pro Ile Pro Asp 315 310 Tyr Met Pro Leu Trp Val Tyr Trp Phe Ile Thr Gly Ile Ser Ile 320 325 Leu Leu Val Gly Ser Val Ile Leu Leu Ile Val Cys Met Thr Trp 335 340 345

```
Arg Leu Ala Gly Pro Gly Ser Glu Lys Tyr Ser Asp Asp Thr Lys
                 350
                                     355
Tyr Thr Asp Gly Leu Pro Ala Ala Asp Leu Ile Pro Pro Pro Leu
                365
                                     370
Lys Pro Arg Lys Val Trp Ile Ile Tyr Ser Ala Asp His Pro Leu
                380
                                     385
Tyr Val Asp Val Val Leu Lys Phe Ala Gln Phe Leu Leu Thr Ala
                 395
                                     400
Cys Gly Thr Glu Val Ala Leu Asp Leu Leu Glu Glu Gln Ala Ile
                 410
                                     415
Ser Glu Ala Gly Val Met Thr Trp Val Gly Arg Gln Lys Gln Glu
                 425
                                     430
Met Val Glu Ser Asn Ser Lys Ile Ile Val Leu Cys Ser Arg Gly
                440
                                     445
                                                          450
Thr Arg Ala Lys Trp Gln Ala Leu Leu Gly Arg Gly Ala Pro Val
                 455
                                     460
Arg Leu Arg Cys Asp His Gly Lys Pro Val Gly Asp Leu Phe Thr
                470
                                     475
Ala Ala Met Asn Met Ile Leu Pro Asp Phe Lys Arg Pro Ala Cys
                                     490
                485
Phe Gly Thr Tyr Val Val Cys Tyr Phe Ser Glu Val Ser Cys Asp
                500
                                     505
Gly Asp Val Pro Asp Leu Phe Gly Ala Ala Pro Arg Tyr Pro Leu
                515
                                     520
Met Asp Arg Phe Glu Glu Val Tyr Phe Arg Ile Gln Asp Leu Glu
                530
                                                          540
                                     535
Met Phe Gln Pro Gly Arg Met His Arg Val Gly Glu Leu Ser Gly
                545
                                     550
                                                          555
Asp Asn Tyr Leu Arg Ser Pro Gly Gly Arg Gln Leu Arg Ala Ala
                560
                                     565
                                                          570
Leu Asp Arg Phe Arg Asp Trp Gln Val Arg Cys Pro Asp Trp Phe
                                     580
                575
                                                          585
Glu Cys Glu Asn Leu Tyr Ser Ala Asp Asp Gln Asp Ala Pro Ser
                590
                                     595
Leu Asp Glu Glu Val Phe Glu Glu Pro Leu Leu Pro Pro Gly Thr
                605
                                     610
Gly Ile Val Lys Arg Ala Pro Leu Val Arg Glu Pro Gly Ser Gln
                620
                                     625
Ala Cys Leu Ala Ile Asp Pro Leu Val Gly Glu Glu Gly Gly Ala
                635
Ala Val Ala Lys Leu Glu Pro His Leu Gln Pro Arg Gly Gln Pro
                650
                                     655
Ala Pro Gln Pro Leu His Thr Leu Val Leu Ala Ala Glu Glu Gly
                665
                                     670
Ala Leu Val Ala Ala Val Glu Pro Gly Pro Leu Ala Asp Gly Ala
                680
                                     685
Ala Val Arg Leu Ala Leu Ala Gly Glu Gly Glu Ala Cys Pro Leu
                                                          705
                695
Leu Gly Ser Pro Gly Ala Gly Arg Asn Ser Val Leu Phe Leu Pro
                710
                                     715
Val Asp Pro Glu Asp Ser Pro Leu Gly Ser Ser Thr Pro Met Ala
                725
                                     730
                                                          735
```

```
Ser Pro Asp Leu Leu Pro Glu Asp Val Arg Glu His Leu Glu Gly
                                                          750
                740
                                     745
Leu Met Leu Ser Leu Phe Glu Gln Ser Leu Ser Cys Gln Ala Gln
                755
                                     760
Gly Gly Cys Ser Arg Pro Ala Met Val Leu Thr Asp Pro His Thr
                                     775
                770
                                                          780
Pro Tyr Glu Glu Glu Gln Arg Gln Ser Val Gln Ser Asp Gln Gly
                785
                                     790
                                                          795
Tyr Ile Ser Arg Ser Ser Pro Gln Pro Pro Glu Gly Leu Thr Glu
                                                          810
                800
                                     805
Met Glu Glu Glu Glu Glu Glu Gln Asp Pro Gly Lys Pro Ala
                815
                                     820
                                                          825
Leu Pro Leu Ser Pro Glu Asp Leu Glu Ser Leu Pro Ser Leu Gln
                                     835
                                                          840
                830
Arg Gln Leu Leu Phe Arg Gln Leu Gln Lys Asn Ser Gly Trp Asp
                                     850
                                                          855
                845
Thr Met Gly Ser Glu Ser Glu Gly Pro Ser Ala
                860
                                     865
```